Amendments to the Specification:

Please note: to distinguish text passages which were already underlined, additions to the text are marked with <u>double underlines</u>.

Please replace the paragraph at page 9, from line 17 through line 30, with the following paragraph:

--The molecule may be a fusion polypeptide which comprises one or more amino acids interposed between the first and second parts which bind to cells, e.g. a fusion polypeptide which comprises a first amino acid sequence which can bind to an antigen bearing target and a second amino acid sequence which can bind to a leukocyte, and which further comprises at least one amino acid interposed between the first and second parts. The interposed amino acids may comprise, e.g., a linker sequence intended to lessen steric hindrance or other undesirable interactions between the aforementioned first and second parts. For, example, one such type of sequence takes the form (Gly₃Ser)_n. Additional useful linkers include, but are not limited to (Arg-Ala-Arg-Asp-Pro-Arg-Val-Pro-Val-Ala-Thr)₁₋₅ (SEQ ID NO:1) (Xu et al., 1999, Proc. Natl. Acad. Sci. U.S.A. 96: 151-156), (Gly-Ser)_n (Shao et al., 2000, Bioconjug. Chem. 11: 822-826), (Thr-Ser-Pro)_n (Kroon et al., 2000, Eur. J. Biochem. 267: 6740-6752), (Gly-Gly-Gly)_n (Kluczyk et al., 2000, Peptides 21: 1411-1420), and (Glu-Lys)_n (Klyczyk et al., 2000, supra), wherein n is 1 to 15 (each of the preceding references is also incorporated herein by reference). In another embodiment, no amino acids are interposed between the first and second parts. –

Please replace the paragraph at page 79, from line 14 through line 20, with the following paragraph:

--At the nucleotide level the human and the murine IL4 gene display approximately 70% homology. The 5' region of the IL4 contains several sequence elements, designated CLE (conserved lymphokine element), that are binding sites for transcription factors controlling the expression of this and other genes. A sequence motif, called P sequence (CGAAAATTTCC; SEQ ID NO:1 SEQ ID NO:2) in the 5' region of the human IL4 gene (positions -79 - - 69) is the binding site for a nuclear factor, called NF(P), mediating the response to T-cell activation signals.--

Please replace the line at page 167, line 10, with the following line: --GTX-5 (SEQ ID NO:3)--

Please replace the line at page 167, line 21, with the following line: --GTX-6 (SEQ ID NO:4)--

Please replace the line at page 168, line 24, with the following line: -- <u>Upstream (SEQ ID NO:5)</u>--

Please replace the line at page 169, line 1, with the following line: -- Downstream (SEQ ID NO:6)--

Please replace the line at page 169, line 24, with the following line: -- <u>Upstream Primer (SEQ ID NO:7)</u>--

Please replace the line at page 169, line 26, with the following line: --Downstream (SEQ ID NO:8) --

Please replace the line at page 171, line 1, with the following line: -- <u>Upsteam (SEQ ID NO:9)</u> --

Please replace the line at page 171, line 3, with the following line: -- Downstream (SEQ ID NO:10) --

Please replace the line at page 178, line 24, with the following line: -- <u>Upstream (SEQ ID NO:11)</u> --

Please replace the line at page 178, line 24, with the following line:
--Downstream (SEO ID NO:12) --

Please replace the paragraph at page 179, from line 25 through line 26, with the following paragraph:

--(SEQ ID NO:13) 5'TACGGCCGGCACCCGCCCGCTCGCCCAGCCCC (SEQ ID NO:14) 3'TACGGCCGCCACAATGAAAATAAGATACCAT --

Please replace the paragraph at page 181, from line 6 through line 7, with the following paragraph:

--<u>(SEQ ID NO:15)</u> 5' CCGGCACTAGTGGCGGAGGGGGCTCCGGCGGGGGGGCAGCG (SEQ ID NO:16) 5' CTAGCGCTGCCCCCGCCGCCGCCGCCCCCCCCCCCACTAGTG --

Please replace the paragraph at page 181, from line 11 through line 14, with the following paragraph:

-- 3. DNA sequence coding for the peptide GGGGSGGGS (SEQ ID NO:17) where G stands for glycine and S stands for serine. This 10 amino acid sequence (G₄S)₂ is designed to insert a kink/spacer in the protein between the GMCSF and the Gas1.1 moieties.--

Please replace the line at page 182, line 14, with the following line:

-- Upstream HA1 Primer (SEQ ID NO:18) --

Please replace the line at page 182, line 16, with the following line:

-- Downstream HA1 Primer (SEQ ID NO:19) --

Please replace the line at page 183, line 9, with the following line:

-- Upstream Primer (SEQ ID NO:20) --

Please replace the line at page 183, line 11, with the following line:
--Downstream Primer (SEQ ID NO:21) --

Please replace the line at page 185, line 26, with the following line:
--Upstream Primer (SEQ ID NO:22) --

Please replace the line at page 185, line 28, with the following line:
--Downstream Primer (SEQ ID NO:23) --

Please replace the line at page 186, line 22, with the following line:
--Upstream Primer (SEQ ID NO:24) --

Please replace the line at page 186, line 24, with the following line:
--Downstream Primer (SEQ ID NO:25) --

Please replace the line at page 187, line 23, with the following line:
--Upstream Primer (SEQ ID NO:26) --

Please replace the line at page 188, line 1, with the following line:
--Downstream Primer (SEO ID NO:27) --

Please replace the line at page 193, line 11, with the following line:
-- Upstream hGM-CSF Primer (SEQ ID NO:28) --

Please replace the line at page 193, line 13, with the following line:
-- Downstream hGM-CSF Primer (SEQ ID NO:29) --

Please replace the line at page 194, line 6, with the following line:
--Upstream Primer (SEQ ID NO:30) --

Please replace the line at page 194, line 8, with the following line:

-- Downstream Primer (SEQ ID NO:31) --

Date: Filmany 24, 2004

Respectfully submitted,

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